| Project Title | Funding | Strategic Plan Objective | Institution | |
|--|-------------|--------------------------|---|--|
| 1/4-The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes | \$817,786 | Q3.S.A | Mount Sinai School of Medicine | |
| 2/4-The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes | \$483,807 | Q3.S.A | Broad Institute, Inc. | |
| An exploration of genetic and behavioral variables in Autism Spectrum Disorder | \$30,800 | Q3.S.A | Center for Autism and Related Disorders (CARD) | |
| Sporadic mutations and autism spectrum disorders | \$713,231 | Q3.S.A | University of Washington | |
| 4/4 The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes | \$759,778 | Q3.S.A | University of California, San Francisco | |
| Autism genetics: Homozygosity mapping and functional validation | \$735,107 | Q3.S.A | Boston Children's Hospital | |
| Genomic influences on development and outcomes in Infants at risk of ASD | \$337,779 | Q3.S.A | University of Alberta | |
| Genomic influences on developmental course and outcome in Infants at risk of ASD: A Baby Siblings Research Consortium (BSRC) Study | \$149,882 | Q3.S.A | University of Alberta | |
| 3/4 - The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes | \$276,478 | Q3.S.A | University of Pittsburgh | |
| Genome-wide expression profiling data analysis to study autism genetic models | \$0 | Q3.S.A | University of California, Los Angeles | |
| Rapid phenotyping for rare variant discovery in autism | \$661,281 | Q3.S.A | University of California, Los Angeles | |
| Environmental exposures measured in deciduous teeth as potential biomarkers for autism risk | \$0 | Q3.S.B | University of Texas Health Science Center at San Antonio | |
| Prenatal and neonatal biologic markers for autism | \$725,197 | Q3.S.C | Kaiser Foundation Research Institute | |
| Air pollution, MET genotype and ASD risk: GxE Interaction in the EMA Study | \$150,000 | Q3.S.C | Kaiser Permanente | |
| The role of germline mutation and parental age in autism spectrum disorders | \$743,939 | Q3.S.C | University of California, San Diego | |
| Perinatal exposure to airborne pollutants and associations with autism phenotype | \$149,737 | Q3.S.C | University of Southern California | |
| The CHARGE study: childhood autism risks from genetics and the environment | \$1,151,250 | Q3.S.C | University of California, Davis | |
| Autism risk, prenatal environmental exposures, and pathophysiologic markers | \$1,759,913 | Q3.S.C | University of California, Davis | |
| ACE Network: Autism Genetics, Phase II: Increasing representation of human diversity | \$3,005,916 | Q3.S.D | University of California, Los Angeles | |
| ACE Network: Autism Genetics, Phase II: Increasing representation of human diversity | \$162,535 | Q3.S.D | University of California, Los Angeles | |
| Prevalence and patterns of medical co-morbidity and healthcare use before ASD diagnoses in children | \$149,999 | Q3.S.E | Kaiser Foundation Research Institute | |
| Maternal autoreactivity and autoimmune disease in autism | \$0 | Q3.S.E | The Feinstein Institute for Medical Research | |
| Research project about a potential infectious origin of autism | \$0 | Q3.S.E | Institut de Recherche Luc Montagnier | |
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| Project Title | Funding | Strategic Plan Objective | Institution | |
|---|-------------|--------------------------|--|--|
| PROTEOMIC MAPPING OF THE IMMUNE RESPONSE TO GLUTEN IN CHILDREN WITH AUTISM | \$67,041 | Q3.S.E | Columbia University New York Morningside | |
| To Study Maternal Anti-GAD Antibodies in Autism | \$5,260 | Q3.S.E | Hartwick College | |
| Environmental exposure unveils mitochondrial dysfunction in autism | \$60,000 | Q3.S.E | University of California, Davis | |
| Novel Proteomics Approach to Oxidative Posttranslational Modifications Underlying Anxiety and Autism Spectrum Disorders | \$0 | Q3.S.E | Sanford Burnham Medical Research Center | |
| UC Davis Center for Children's Environmental Health (CCEH) Bridge | \$0 | Q3.S.F | University of California, Davis | |
| Epidemiological research on autism in Jamaica - Phase II | \$607,366 | Q3.S.H | University of Texas Health Science Center at Houston | |
| Parental Exposures to Occupational Asthmagens and Risk of Autism Spectrum Disorders | \$29,500 | Q3.S.H | Johns Hopkins University | |
| Gestational exposure questionnaire validation and feasibility study | \$20,262 | Q3.S.H | University of California, Davis | |
| Gestational metabolic conditions and autism | \$77,000 | Q3.S.H | University of California, Davis | |
| Very early behavioral indicators of ASD risk among NICU infants: A prospective study | \$149,986 | Q3.S.H | Institute for Basic Research in Developmental Disabilities | |
| Neonatal biomarkers in extremely preterm babies predict childhood brain disorders | \$3,655,744 | Q3.S.H | Boston Medical Center | |
| Early life environmental exposures and autism in an existing Swedish birth cohort | \$0 | Q3.S.H | Drexel University | |
| Risk factors, comorbid conditions, and epidemiology of autism in children | \$0 | Q3.S.H | Henry M. Jackson Foundation | |
| Prenatal folic acid and risk for autism spectrum disorders | \$124,870 | Q3.S.H | Emory University School of Medicine | |
| Investigation of Transgenerational Neurodevelopmental Impacts of Gestational Pharmaceuticals | \$100,000 | Q3.S.H | Institute of Preventive Medicine at Frederiksberg Hospital | |
| Prenatal Androgen in Meconium and Early Autism Spectrum Disorder Related Neurodevelopmental Outcomes | \$29,423 | Q3.S.H | Drexel University | |
| In utero antidepressant exposures and risk for autism | \$343,560 | Q3.S.H | Massachusetts General Hospital | |
| Defining the underlying biology of gastrointestinal dysfunction in autism | \$0 | Q3.S.I | University of California, Davis | |
| Modeling gut microbial ecology and metabolism in autism using an innovative ex vivo approach | \$0 | Q3.S.I | University of Guelph | |
| Modeling Gut Microbial Ecology and Metabolism in Autism Using an Innovative Ex Vivo Approach | \$22,441 | Q3.S.I | University of Guelph | |
| Autism, GI symptoms and the enteric microbiota | \$350,814 | Q3.S.I | The Research Foundation of the State University of New York at Stony Brook | |
| Investigating the gut microbiome for novel therapies and diagnostics for autism | \$558,136 | Q3.S.I | California Institute of Technology | |

| Project Title | Funding | Strategic Plan Objective | Institution | |
|---|-------------|--------------------------|---|--|
| Elevated urinary P-cresol in small autistic children: Origin and consequences | \$0 | Q3.S.I | Universita Campus Bio-Medico di Roma | |
| Regressive autism as an infectious disease: Role of the home as an environmental factor | \$0 | Q3.S.I | VA Medical Center, Los Angeles | |
| Role of Intestinal Microbiome in Children with Autism | \$29,000 | Q3.S.I | Massachusetts General Hospital | |
| Human neurobehavioral phenotypes associates with the extended PWS/AS domain | \$587,398 | Q3.S.J | Baylor College of Medicine | |
| Genome-wide analyses of DNA methylation in autism | \$0 | Q3.S.J | Mount Sinai School of Medicine | |
| 5-Hydroxymethylcytocine-mediated epigenetic regulation in autism spectrum disorders | \$60,000 | Q3.S.J | Emory University | |
| 5-hydroxymethylcytocine-mediated epigenetic regulation in autism | \$200,000 | Q3.S.J | Emory University | |
| In vivo function of neuronal activity-induced MeCP2 phosphorylation | \$277,792 | Q3.S.J | University of Wisconsin - Madison | |
| Epigenetic DNA modifications in autistic spectrum disorders | \$163,813 | Q3.S.J | Johns Hopkins University School of Medicine | |
| Mechanisms of valproic acid-induced neurodevelopmental and behavioral defects | \$302,269 | Q3.S.J | University of Maryland, Baltimore | |
| Epigenetic and transcriptional dysregulation in autism spectrum disorder | \$748,775 | Q3.S.J | University of California, Los Angeles | |
| Methylomic and genomic impacts of organic pollutants in Dup15q syndrome | \$338,560 | Q3.S.J | University of California, Davis | |
| Project 2: Perinatal epigenetic signature of environmental exposure | \$105,416 | Q3.S.J | University of California, Davis | |
| Genome-wide examination of DNA methylation in autism | \$149,999 | Q3.S.J | Johns Hopkins University | |
| Regulation of gene expression through complex containing AUTS2 | \$100,854 | Q3.S.J | New York University School of Medicine | |
| Exploring interactions between folate and environmental risk factors for autism | \$153,615 | Q3.S.J | University of California, Davis | |
| Mutations in heterochromatin-related genes in autism | \$0 | Q3.S.J | Hebrew University of Jerusalem | |
| Cell specific genomic imprinfing during cortical development and in mouse models | \$308,216 | Q3.S.J | Harvard University | |
| Environment, the perinatal epigenome, and risk for autism and related disorders | \$1,400,550 | Q3.S.J | Johns Hopkins University | |
| Paternal age and epigenetic mechanisms in psychiatric disease | \$15,000 | Q3.S.J | Research Foundation for Mental Hygiene, Inc/NYSPI | |
| Evaluating the Functional Impact of Epigenetic Control Related Genes Mutated in both Schizophrenia and Autism | \$0 | Q3.S.J | Columbia University | |
| Conservation of imprinting for autism-linked genes in the brain | \$60,000 | Q3.S.J | University of Utah | |

| Project Title | Funding | Strategic Plan Objective | Institution |
|---|-------------|--------------------------|---|
| Genetic and environmental interactions leading to autism-like symptoms | \$0 | Q3.S.K | The Rockefeller University |
| The role of serotonin in social bonding in animal models | \$0 | Q3.S.K | University of California, Davis |
| Cellular and Synaptic Dissection of the Neuronal Circuits of Social and Autistic Behavior | \$0 | Q3.S.K | University of Coimbra |
| Evaluating epidemiological and biostatistical challenges n the EARLI investigation | \$0 | Q3.L.A | Drexel University |
| Early autism risk longitudinal investigation (EARLI) network | \$411,571 | Q3.L.A | Drexel University |
| ntegrative genetic analysis of autism brain tissue | \$0 | Q3.L.B | Johns Hopkins University School of Medicine |
| imons Simplex Collection support grant | \$24,484 | Q3.L.B | University of Washington |
| Sequencing Female-enriched Multiplex Autism Families FEMFs) | \$0 | Q3.L.B | Johns Hopkins University School of Medicine |
| Simons Simplex Collection support grant | \$23,645 | Q3.L.B | University of Illinois at Chicago |
| /3-Sequencing autism spectrum disorder extended pedigrees | \$286,240 | Q3.L.B | University of Utah |
| Genomic hotspots of autism | \$0 | Q3.L.B | University of Washington |
| Whole exome sequencing of Simons Simplex Collection uads | \$1,495,957 | Q3.L.B | University of Washington |
| Senetic basis of phenotypic variability in 16p11.2 leletion or duplication | \$0 | Q3.L.B | University of Washington |
| genome-wide search for autism genes in the SSC imory | \$0 | Q3.L.B | Emory University |
| dypocholesterolemic autism spectrum disorder | \$45,647 | Q3.L.B | National Institutes of Health |
| /3-Sequencing autism spectrum disorder extended edigrees | \$153,600 | Q3.L.B | University of Pennsylvania |
| autism Genome Project (AGP): Genome sequencing and analysis supplement | \$0 | Q3.L.B | The Hospital for Sick Children |
| examining the Y-chromosome in autism spectrum isorder | \$0 | Q3.L.B | The Hospital for Sick Children |
| Mutations in noncoding DNA and the missing heritability f autism | \$124,987 | Q3.L.B | University of California, San Diego |
| nvestigation of DUF1220 domains in human brain unction and disease | \$361,544 | Q3.L.B | University of Colorado Denver |
| Whole exome sequencing of Simons Simplex Collection uads | \$536,779 | Q3.L.B | Yale University |
| imons Simplex Collection support grant | \$25,735 | Q3.L.B | Vanderbilt University Medical Center |
| genome-wide search for autism genes in the SSC | \$0 | Q3.L.B | Vanderbilt University Medical Center |

| Project Title | Funding | Strategic Plan Objective | Institution |
|--|-------------|--------------------------|--|
| Complex genetic architecture of chromosomal aberrations in autism | \$92,917 | Q3.L.B | Massachusetts General Hospital |
| Cryptic chromosomal aberrations contributing to autism | \$135,649 | Q3.L.B | Massachusetts General Hospital |
| Genetic epidemiology of complex traits | \$589,154 | Q3.L.B | National Institutes of Health |
| Mitochondria and the etiology of autism | \$350,000 | Q3.L.B | The Children's Hospital of Philadelphia |
| Autism genetics: Homozygosity mapping and functional validation | \$150,000 | Q3.L.B | Boston Children's Hospital |
| Simons Simplex Collection support grant | \$23,171 | Q3.L.B | Boston Children's Hospital |
| Finding recessive genes for autism spectrum disorders | \$175,000 | Q3.L.B | Boston Children's Hospital |
| Genetic basis of autism | \$4,000,571 | Q3.L.B | Cold Spring Harbor Laboratory |
| 2/3-Sequencing autism spectrum disorder extended pedigrees | \$222,480 | Q3.L.B | University of Washington |
| Whole-exome sequencing to identify causative genes for autism | \$175,000 | Q3.L.B | Rockefeller University |
| Dissecting expression regulation of an autism GWAS hit | \$15,000 | Q3.L.B | University of California, San Francisco |
| Genomic influences on development and outcomes in infants at risk for autism | \$681,108 | Q3.L.B | University of Alberta |
| Genomic profiling of autism families using whole- genome sequencing | \$174,960 | Q3.L.B | Institut Pasteur |
| Novel statistical methods for DNA sequencing data, and applications to autism | \$314,312 | Q3.L.B | Columbia University |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | University of California, Los Angeles |
| A genome-wide search for autism genes in the SSC UCLA | \$0 | Q3.L.B | University of California, Los Angeles |
| Simons Simplex Collection support grant | \$26,824 | Q3.L.B | Baylor College of Medicine |
| Simons Simplex Collection support grant | \$20,991 | Q3.L.B | Weill Cornell Medical College |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | University of Missouri |
| llumina, Inc. | \$556,250 | Q3.L.B | Illumina, Inc. |
| Autism Genome Project (AGP) | \$0 | Q3.L.B | Autism Speaks (AS) |
| Simons Simplex Collection support grant | \$25,704 | Q3.L.B | Yale University |
| Simons Simplex Collection support grant | \$21,675 | Q3.L.B | Columbia University |
| Simons Simplex Collection support grant | \$30,000 | Q3.L.B | Emory University |
| Sequence-based discovery of genes with pleiotropic effects across diagnostic boundaries and throughout the ifespan | \$29,995 | Q3.L.B | Massachusetts General Hospital and Harvard Universit |
| Autism Genome Project Consortium data reanalysis using computational biostatistics | \$0 | Q3.L.B | The Rockefeller University |

| Project Title | Funding | Strategic Plan Objective | Institution | |
|---|-------------|--------------------------|---|--|
| Identifying genetic variants on the Y chromosome of males with autism | \$53,430 | Q3.L.B | The Hospital for Sick Children | |
| Next generation gene discovery in familial autism | \$644,126 | Q3.L.B | University of Washington | |
| Identification of functional networks perturbed in autism | \$60,000 | Q3.L.B | Columbia University | |
| Developing new statisical methods to detect variants involved in complex disease | \$434,485 | Q3.L.B | National Institutes of Health | |
| Molecular Characterization of Autism Gene CHD8 in Shaping the Brain Epigenome | \$35,000 | Q3.L.B | Boston Children's Hospital | |
| Genome-wide analysis of cis-regulatory elements in autism | \$62,500 | Q3.L.B | Washington University in St. Louis | |
| Simons Simplex Collection support grant | \$21,268 | Q3.L.B | McGill University Health Centre- Montreal Children's Hospital | |
| PLACENTAL IDENTIFICATION AND IMMUNE QUANTIFICATION OF ACUTE AND/OR CHRONIC INFLAMMATION IN CHILDREN DIAGNOSED WITH PLACENTAL AUTISM IN UNIVERSITY AND COMMUNITY HOSPITALS | \$148,000 | Q3.L.C | Institute for Basic Research in Developmental Disabilities | |
| Prenatal antidepressants and autism spectrum disorder | \$0 | Q3.L.C | Cincinnati Children's Hospital Medical Center | |
| Prenatal PBDE exposure and ASD-related developmental outcomes in the EARLI cohort | \$150,000 | Q3.L.C | Drexel University | |
| Assisted reproductive technologies and increased autism risk | \$192,000 | Q3.L.C | Columbia University | |
| Centers for Autism and Developmental Disabilities Research and Epidemiology (CADDRE) - Georgia | \$985,604 | Q3.L.D | Centers for Disease Control and Prevention (CDC) | |
| Gene-environment interactions in an autism birth cohort | \$6,537,537 | Q3.L.D | Columbia University | |
| Centers for Autism and Developmental Disabilities Research and Epidemiology (CADDRE) - North Carolina | \$1,050,000 | Q3.L.D | University of North Carolina at Chapel Hill | |
| Centers for Autism and Developmental Disabilities Research and Epidemiology (CADDRE) - California | \$1,050,000 | Q3.L.D | Kaiser Foundation Research Institute | |
| Centers for Autism and Developmental Disabilities Research and Epidemiology (CADDRE) - Colorado | \$1,050,000 | Q3.L.D | Colorado Department of Health and Environment | |
| Population-based autism genetics & environment study | \$600,532 | Q3.L.D | Mount Sinai School of Medicine | |
| Parental age and schizophrenia susceptibility | \$308,000 | Q3.L.D | University of California, Los Angeles | |
| The UC Davis Center for Children's Environmental Health and Disease Prevention | \$1,660,178 | Q3.L.D | University of California - Davis | |
| Community-based study of autism spectrum disorders among 7-9 y old children in rural Bangladesh | \$0 | Q3.L.D | Johns Hopkins University | |
| Centers for Autism and Developmental Disabilities Research and Epidemiology (CADDRE) - Maryland | \$1,000,000 | Q3.L.D | Johns Hopkins University | |
| Project 1: Epidemiology and the environment in autism (Hertz-Picciotto) | \$158,613 | Q3.L.D | University of California, Davis | |

| Project Title | Funding | Strategic Plan Objective | Institution |
|---|-------------|--------------------------|--|
| The roles of environmental risks and GEX in increasing ASD prevalence | \$532,325 | Q3.L.D | Yale University |
| Centers for Autism and Developmental Disabilities Research and Epidemiology (CADDRE) - Pennsylvania | \$1,050,000 | Q3.L.D | University of Pennsylvania/Children's Hospital of Philadelphia |
| Centers for Autism and Developmental Disabilities Research and Epidemiology (CADDRE) - Data Coordinating Center | \$868,500 | Q3.L.D | Michigan State University |
| ACE Network: Multigenerational Familial and Environmental Risk for Autism (MINERvA) Network | \$948,404 | Q3.L.D | Mount Sinai School of Medicine |
| FOXP2-regulated signaling pathways critical for higher cognitive functions | \$291,826 | Q3.Other | University of Texas Southwestern Medical Center |
| A history of behavioral genetics | \$0 | Q3.Other | University of Pittsburgh |
| Non-coding RNAs in autism | \$246,000 | Q3.Other | University of Southern California |